## IN THE CLAIMS:

Please amend claims 1-6, 8-13, and add new claims 14-15 as follows:

(Currently Amended) A method for displaying a dendrogram comprising the steps of:
clustering a plurality [[of]] types of biopolymers based on a set of data obtained
by experiments of the plurality types of biopolymers under different conditions, and
displaying [[the]] clustering results thereof in a form of a dendrogram format;

selecting a subtree in the dendrogram; and

displaying the selected subtree on a separate window thereby grouping biopolymers in the selected subtree into at least one function unit or function group.

2. (Currently Amended) A method for displaying a dendrogram according to claim 1, further comprising the steps of:

designating a different clustering method for [[the]] biopolymers included in the subtree displayed on the separate window; and

secondarily clustering the biopolymers included in the subtree [[again]] according to the designated clustering method, and displaying [[the]] secondarily clustering results thereof in a form of a dendrogram format.

3. (Currently Amended) A method for displaying a dendrogram comprising the steps of:

clustering a plurality [[of]] types of biopolymers based on a set of data obtained by experiments of the plurality <u>types</u> of biopolymers under different conditions, and displaying [[the]] <u>clustering</u> results thereof in a form of a dendrogram <del>format</del>;

selecting a subtree in the dendrogram; [[and]]

replacing the selected subtree with an icon in the dendrogram thereby grouping biopolymers in the selected subtree into at least one function unit or function group.

4. (Currently Amended) A method for displaying a dendrogram according to claim 3, further comprising a step of restoring the subtree icon <u>back</u> to the original dendrogram subtree in the dendrogram format.

(Currently Amended) A method for displaying a dendrogram comprising the steps of: clustering a plurality [[of]] types of biopolymers based on a set of data obtained by experiments of the plurality types of biopolymers under different conditions, and displaying [[the]] clustering results thereof in a form of a dendrogram format;

selecting a subtree in the dendrogram; and

from the biopolymers included in the selected subtree, counting and displaying predetermined keywords and a corresponding [[the]] number of biopolymers containing in [[their]] biopolymer information thereof a respective one of the predetermined keywords thereby grouping biopolymers in the selected subtree into at least one function unit or function group from a keyword dictionary file.

6. (Currently Amended) A method for displaying a dendrogram comprising the steps of: clustering a plurality [[of]] types of biopolymers based on a set of data obtained by experiments of the plurality types of biopolymers under different conditions, and

displaying [[the]] clustering results thereof in a form of a dendrogram format;

selecting a subtree in the dendrogram;

designating [[a]] at least one keyword for the selected subtree; and

displaying the selected subtree and highlighting a location of [[a]] each biopolymer in the selected subtree dendrogram, which includes the designated keyword in [[its]] biopolymer information thereof thereby grouping biopolymers in the selected subtree into at least one function unit or function group.

- 7. (Original) A method for displaying a dendrogram according to any one of claims 1 to 6, wherein the biopolymers are cDNAs, RNAs, DNA fragments or genes.
- 8. (Currently Amended) A system for displaying a dendrogram comprising:

a clustering processor for clustering a plurality [[of]] types of biopolymers based on a set of data obtained by experiments of the plurality types of biopolymers under different conditions, and analyzing [[the]] and displaying clustering results thereof to display them in a form of a dendrogram format;

a display system for displaying the dendrogram and for displaying on a separate

window a subtree selected by a user thereby grouping biopolymers in the selected subtree into at least one function unit or function group; and

- a keyword dictionary file for storing keywords of biopolymer information associated with each of the plurality types of biopolymers.
- 9. (Currently Amended) A system for displaying a dendrogram according to claim 8, <u>further</u> comprising a <u>function of displaying a subtree selected by the</u> input means <u>for selecting</u> the subtree by the user on a separate window.
- 10. (Currently Amended) A system for displaying a dendrogram according to claim [[9]] 8, further comprising a function of means for designating a different clustering method for the subtree displayed on the separate window to secondarily cluster the biopolymers included in the subtree again according to the designated clustering method, and displaying [[the]] secondarily clustering results thereof in a form of a dendrogram format.
- 11. (Currently Amended) A system for displaying a dendrogram according to any one of claims 8 to 10, wherein the system comprises a function of further comprising means for replacing the selected subtree selected by the input means with an icon, and a function of means for restoring the subtree icon back to the original subtree in the dendrogram format..
- 12. (Currently Amended) A system for displaying a dendrogram according to any one of claims 8 to [[11]]10, wherein the system comprises a function of further comprising one of means for counting and displaying predetermined keywords retrieved from the keyword dictionary file and a corresponding [[the]] number of biopolymers containing in [[their]] biopolymer information thereof a respective one of the predetermined keywords from a keyword dictionary file, and/or a function of displaying highlighting a location of [[a]] each biopolymer in the selected subtree dendrogram, which includes the designated predetermined keywords in the biopolymer information thereof.

- 13. (Currently Amended) A system for displaying a dendrogram according to any one of claims 8 to [[12]]10, wherein the biopolymers are cDNAs, RNAs, DNA fragments or genes.
- 14. (New) A method for displaying a dendrogram according to claim 5, wherein the counting step involves counting synonyms of the respective one of the predetermined keywords.
- 15. (New) A system for displaying a dendrogram according to claim 12, wherein the means for counting and displaying counts synonyms of each of the predetermined keywords.